bio-protocol Preprint

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2P-seq

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An abbreviated version of this protocol was published in eLIFE in Mar 2018

Codon usage biases co-evolve with transcription termination machinery to suppress premature cleavage and polyadenylation

DOI: 10.7554/eLife.33569

Related files

2p-seq protocol.pdf

How to cite:(Readers should cite both the Bio-protocol preprint and the original research article where this protocol was used)

1. Dang, Y. , Liu, Y. and Zhou, Z. (2020). 2P-seq. Bio-protocol Preprint. bio-protocol.org/prep596.
2. Zhou, Z., Dang, Y., Zhou, M., Yuan, H. and Liu, Y.(2018). Codon usage biases co-evolve with transcription termination machinery to suppress premature cleavage and polyadenylation. eLIFE. DOI: 10.7554/eLife.33569

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